

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 1015801176
Source: JPL/DO
Date Processed by STIC: 4/20/07

ENTERED



IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/580,176

DATE: 04/20/2007

TIME: 16:03:59

Input Set : E:\sequence listing yeda-039 ST25.txt
 Output Set: N:\CRF4\04202007\J580176.raw

3 <110> APPLICANT: YEDA Research and Development
 5 <120> TITLE OF INVENTION: DNA VACCINES ENCODING HSP60 PEPTIDE FRAGMENTS FOR TREATING
 6 AUTOIMMUNE DISEASES
 8 <130> FILE REFERENCE: YEDA/039 PCT
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/580,176
 C--> 10 <141> CURRENT FILING DATE: 2006-05-22
 10 <160> NUMBER OF SEQ ID NOS: 15
 12 <170> SOFTWARE: PatentIn version 3.3
 14 <210> SEQ ID NO: 1
 15 <211> LENGTH: 20
 16 <212> TYPE: PRT
 17 <213> ORGANISM: Artificial
 19 <220> FEATURE:
 20 <223> OTHER INFORMATION: Human HSP60 epitope
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 28 Val Leu Asn Arg
 29 20
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 52 <212> TYPE: PRT
 53 <213> ORGANISM: Artificial
 55 <220> FEATURE:
 56 <223> OTHER INFORMATION: Human HSP60 epitope
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 64 Gln Leu Asp Val
 65 20
 68 <210> SEQ ID NO: 4
 69 <211> LENGTH: 20

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73 <220> FEATURE:
74 <223> OTHER INFORMATION: Human HSP60 epitope
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89 <213> ORGANISM: Artificial
91 <220> FEATURE:
92 <223> OTHER INFORMATION: Human HSP60 epitope
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125 <213> ORGANISM: Artificial
127 <220> FEATURE:
128 <223> OTHER INFORMATION: Human HSP60 epitope
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206 <211> LENGTH: 2691
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208 <213> ORGANISM: Homo sapiens
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215 tgattggtcc aaggaaggct gggggcagg acgggaggcg aaaccctgg aatattcccg 180
217 acctggcagc ctcatcgagc tcggtgattt gctcagaagg gaaaaggcgg gtctccgtga 240
219 cgacttataaa aagcccgagg gcaagcggtc cggataacgg cttagcctgag gagctgctgc 300
221 gacagtccac tactttttc gagagtact cccgttgtcc caaggcttcc cagagcgaac 360
223 ctgtgcggct gcaggccaccg gcgcgtcgag tttccggcgt ccggaaaggac cgagctttc 420
225 tcgcggatcc agtgtccgt ttccagcccc caatctcaga gccgagccga cagagagcag 480
227 ggaaccgcatt gccaacaaagcc gcggcagtcg gcatcgacct gggcaccacc tactcctgcg 540
229 tgggggtgtt ccaacacggc aagggtggaga tcatcgccaa cgaccaggc aaccgcacca 600
231 ccccccagcta cgtggccttc acggacacccg agcggctcat cggggatgcg gccaagaacc 660

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233	aggtggcgct	gaaccgcag	aacaccgtgt	ttgacgcgaa	gcgcctgatc	ggccgcaagt	720
235	tcggcgaccc	ggtggtcag	tcggacatga	agcactggcc	tttccaggtg	atcaacgacg	780
237	gagacaagcc	caagggtcag	gtgagctaca	agggggagac	caaggcattc	taccccgagg	840
239	agatctcgtc	catggtgctg	accaagatga	aggagatcgc	cgaggcgta	ctgggctacc	900
241	cggtgaccaa	cgcggtgatc	accgtgccgg	cctacttcaa	cgactcgcag	cgccaggcca	960
243	ccaaggatgc	gggtgtgatc	gcggggctca	acgtgctgcg	gatcatcaac	gagcccacgg	1020
245	ccgcccacat	cgcctacggc	ctggacagaa	cgggcaaggg	ggagcgcac	gtcctgatct	1080
247	ttgacctggg	cgggggcacc	ttcgacgtgt	ccatcctgac	gatcgacgac	ggcatcttcg	1140
249	aggtgaaggc	cacggccggg	gacacccacc	tgggtgggaa	ggactttgac	aacaggctgg	1200
251	tgaaccactt	cgtggaggag	ttcaagagaa	aacacaagaa	ggacatcgc	cagaacaagc	1260
253	gagccgtgag	gcggctgcgc	accgcctgcg	agagggccaa	gaggaccctg	tcgtccagca	1320
255	cccaggccag	cctggagatc	gactccctgt	ttgagggcat	cgacttctac	acgtccatca	1380
257	ccagggcggag	tttcgaggag	ctgtgtcccg	acctgttccg	aagcacccctg	gagcccggtgg	1440
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267	ccctgtcgct	ggggctggag	acggccggag	qcggtatgac	tgcctgatc	aagcgcact	1740
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271	tgctgatcca	ggtgtacgag	ggcgagaggg	ccatgacgaa	agacaacaat	ctgttggggc	1860
273	gttcgagct	gagcggcatc	cctccggccc	caggcgtgcc	ccagatcgcg	gtgaccttcg	1920
275	acatcgatgc	caacggcatc	ctgaacgtca	cggccacgg	caagagcacc	ggcaaggcca	1980
277	acaagatcac	catcaccaac	gacaagggcc	gcctgagcaa	ggaggagatc	gagcgcattgg	2040
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285	cgtggctgga	cgccaaacacc	ttggccgaga	aggacgagtt	tgagcacaag	aggaaggagc	2280
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312	gcctttcagg	cagaaatttc	ccagttgtat	tcattgtatc	tcaataactt	ctactcgaac	180
314	aaagagatct	ttctgagaga	gctcatttca	aattcatcg	atgcatttgg	caaaatccgg	240
316	tatgaaactt	tgacagatcc	cagtaaatta	gactctggg	aagagctgca	tattaacctt	300
318	ataccgaaca	aacaagatcg	aactctca	attgtggata	ctggaaattgg	aatgaccaag	360
320	gctgacttga	tcaataaacct	ttgtactatc	gccaaagtctg	ggaccaaagc	gttcatggaa	420
322	gctttgcagg	ctgggtcaga	tatctctatg	attggccagt	tcgggttgg	tttttattct	480
324	gcttatttgg	ttgctgagaa	agtaactgtg	atcaccaaac	ataacgatga	tgagcgtac	540
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332	tttgtggaga	aggaacgtga	taaagaagta	agcgatgatg	aggctgaaga	aaaggaagac	780										
334	aaagaagaag	aaaaagaaaa	agaagagaaa	gagtcggaag	acaaacctga	aattgaagat	840										
336	gttggttctg	atgaggaaga	agaaaagaag	gatggtgaca	agaagaagaa	gaagaagatt	900										
338	aaggaaaagt	acatcgatca	agaagagctc	aacaaaacaa	agcccatctg	gaccagaaat	960										
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342	gaagatca	tggcagtgaa	gcattttca	gttgaaggac	agtttggatt	cagagccctt	1080										
344	ctatttgc	cacgacgtgc	tcctttgtat	ctgtttgaaa	acagaaaagaa	aaagaacaat	1140										
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378	ccccagacac	atgctaacag	gatctacagg	atgatcaa	tgggtctggg	tattgtgaa	2160										
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419						20				25					30		
422	Gly	Ala	Asp	Ala	Arg	Ala	Leu	Met	Leu	Gln	Gly	Val	Asp	Leu	Leu	Ala	

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Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10

VERIFICATION SUMMARY

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L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date